

WEST Search History

DATE: Monday, March 10, 2003

Set Name Query
side by side

Hit Count Set Name
result set

DB=USPT,PGPB; PLUR=YES; OP=ADJ

L3	(ebna1 or ebna-1 or bkrfl or brkf-1) and (barfl or barf-1)	3	L3
L2	L1 same (primer or probe)	38	L2
L1	(ebna1 or ebna-1) or (barfl or barf1) or (bkrfl or brkf-1)	431	L1

END OF SEARCH HISTORY

Proteins,specific or class...

LMP-2 (latent-infection membrane protein 2), detection of gene for;
primers for amplification and detection of Epstein Barr virus nucleic
acid by NASBA

Gene,microbial...

LMP1, detection of; primers for amplification and detection of Epstein
Barr virus nucleic acid by NASBA

Gene,microbial...

LMP2, detection of; primers for amplification and detection of Epstein
Barr virus nucleic acid by NASBA

Antigens...

LYDMA, detection of gene for; primers for amplification and detection
of Epstein Barr virus nucleic acid by NASBA

Diagnosis...

mol., of Epstein-Barr virus infection; primers for amplification and
detection of Epstein Barr virus nucleic acid by NASBA

Human herpesvirus 4... NASBA(nucleic acid sequence-based amplification)...

Primers(nucleic acid)...

primers for amplification and detection of Epstein Barr virus nucleic
acid by NASBA

Interleukin 10...

·viral, detection of gene for; primers for amplification and detection
of Epstein Barr virus nucleic acid by NASBA

CAS REGISTRY NUMBERS:

219991-34-1 219991-38-5 219991-42-1 219991-43-2 242135-85-9 primer for
detection BARF-1 gene of Epstein Barr virus; primers for amplification
and detection of Epstein Barr virus nucleic acid by NASBA

242135-86-0 242135-87-1 242135-88-2 242135-89-3 242135-99-5 primer for
detection BCRF-1 gene of Epstein Barr virus; primers for amplification
and detection of Epstein Barr virus nucleic acid by NASBA

242135-90-6 242135-91-7 242135-92-8 242135-93-9 242136-00-1 primer for
detection BDLF-2 gene of Epstein Barr virus; primers for amplification
and detection of Epstein Barr virus nucleic acid by NASBA

219990-89-3 219990-97-3 242135-80-4 242135-81-5 242135-82-6
242135-83-7 primer for detection EBER-1 RNA of Epstein Barr virus;
primers for amplification and detection of Epstein Barr virus nucleic
acid by NASBA

219990-30-4 219990-43-9 219990-78-0 242135-72-4 242135-79-1 primer for
detection EBNA-1 gene of Epstein Barr virus; primers for amplification
and detection of Epstein Barr virus nucleic acid by NASBA

219990-99-5 219991-00-1 219991-05-6 219991-13-6 242135-84-8 primer for
detection LMP-1 gene of Epstein Barr virus; primers for amplification
and detection of Epstein Barr virus nucleic acid by NASBA

219991-18-1 219991-25-0 219991-31-8 219991-32-9 219991-33-0 primer for
detection LMP-2 gene of Epstein Barr virus; primers for amplification
and detection of Epstein Barr virus nucleic acid by NASBA

? s (barf1 or barf-1) and (epithelial or breast or nasopharyngeal or npc)

147 BARF1

0 BARF-1

746772 EPITHELIAL

824632 BREAST

43828 NASOPHARYNGEAL

13149 NPC

S3 83 (BARF1 OR BARF-1) AND (EPITHELIAL OR BREAST OR
NASOPHARYNGEAL OR NPC)

? rd s3

...examined 50 records (50)

...completed examining records

S4 22 RD S3 (unique items)

? t s4/6/1-22

4/6/1 (Item 1 from file: 5)

13534988 BIOSIS NO.: 200200163809

Hypothesis: A novel route for immortalization of **epithelial** cells by Epstein-Barr virus.

2002

4/6/2 (Item 2 from file: 5)

13310250 BIOSIS NO.: 200100517399

Growth transformation of primary **epithelial** cells with a NPC -derived Epstein-Barr virus strain.

2001

4/6/3 (Item 3 from file: 5)

12965681 BIOSIS NO.: 200100172830

N-terminal domain of **BARF1** gene encoded by Epstein-Barr virus is essential for malignant transformation of rodent fibroblasts and activation of Bcl-2.

2001

4/6/4 (Item 4 from file: 5)

12748457 BIOSIS NO.: 200000502080

Expression of **BARF1** gene encoded by Epstein-Barr virus in **nasopharyngeal** carcinoma biopsies.

2000

4/6/5 (Item 5 from file: 5)

12528497 BIOSIS NO.: 200000281999

Role of the Epstein-Barr virus Rta protein in activation of distinct classes of viral lytic cycle genes.

1999

4/6/6 (Item 6 from file: 5)

12522429 BIOSIS NO.: 200000275931

Unique transcription pattern of Epstein-Barr virus (EBV) in EBV-carrying gastric adenocarcinomas: Expression of the transforming **BARF1** gene.

2000

4/6/7 (Item 7 from file: 5)

11973916 BIOSIS NO.: 199900227229

Expression of Epstein-Barr virus (EBV) transcripts encoding homologues to important human proteins in diverse EBV associated diseases.

1999

4/6/8 (Item 8 from file: 5)

11708757 BIOSIS NO.: 199800490488

Nucleic acid sequence-based amplification, a new method for analysis of spliced and unspliced Epstein-Barr virus latent transcripts, and its comparison with reverse transcriptase PCR.

1998

4/6/9 (Item 9 from file: 5)

11295044 BIOSIS NO.: 199800076376

Expression of the protein encoded by Epstein-Barr virus (EBV) **BARF1** open reading frame from a recombinant adenovirus system.

1997

4/6/10 (Item 10 from file: 5)
11025142 BIOSIS NO.: 199799646287
Establishment of a monkey kidney **epithelial** cell line with the
BARF1 open reading frame from Epstein-Barr virus.
1997

4/6/11 (Item 11 from file: 5)
10745514 BIOSIS NO.: 199799366659
Antibody and antibody-dependent cellular cytotoxicity responses against the
BamHI A rightward open-reading frame-1 protein of Epstein-Barr virus
(EBV) in EBV-associated disorders.
1997

4/6/12 (Item 1 from file: 34)
08668991 Genuine Article#: 314MJ Number of References: 34
Title: Unique transcription pattern of Epstein-Barr virus (EBV) in
EBV-carrying gastric adenocarcinomas: Expression of the transforming
BARF1 gene (ABSTRACT AVAILABLE).
Publication date: 20000515

4/6/13 (Item 2 from file: 34)
06651219 Genuine Article#: ZH268 Number of References: 60
Title: The Epstein-Barr virus **BARF1** gene encodes a novel, soluble
colony-stimulating factor-1 receptor (ABSTRACT AVAILABLE)
Publication date: 19980500

4/6/14 (Item 3 from file: 34)
03058166 Genuine Article#: ND238 Number of References: 31
Title: EXPRESSION AND TUMORIGENICITY OF THE EPSTEIN-BARR-VIRUS **BARF1**
GENE IN HUMAN LOUCKES B-LYMPHOCYTE CELL-LINE (Abstract Available)

4/6/15 (Item 4 from file: 34)
02084998 Genuine Article#: JZ811 Number of References: 24
Title: TRANSCRIPTIONAL EXPRESSION OF THE VIRAL GENOME IN THE EPSTEIN-BARR
VIRUS-INDUCED TAMARIN LYMPHOMA AND THE CORRESPONDING LYMPHOBLASTOID
TUMOR LINES (Abstract Available)

4/6/16 (Item 1 from file: 76)
02492243 4737903
Unique Transcription Pattern of Epstein-Barr Virus (EBV) in EBV-carrying
Gastric Adenocarcinomas: Expression of the Transforming **BARF1** Gene
(2000)

4/6/17 (Item 2 from file: 76)
02217680 4231116
Expression of the protein encoded by Epstein-Barr virus (EBV) **BARF1**
open reading frame from a recombinant adenovirus system
(1997)

4/6/18 (Item 1 from file: 98)
04273985 H.W. WILSON RECORD NUMBER: BGSA00023985 (USE FORMAT 7 FOR
FULLTEXT)

Role of cytotoxic T lymphocytes in Epstein-Barr virus-associated diseases.
WORD COUNT: 11989
2000 (200000000)

4/6/19 (Item 1 from file: 144)
15110549 PASCAL No.: 01-0271038
N-terminal domain of **BARF1** gene encoded by Epstein-Barr virus is
essential for malignant transformation of rodent fibroblasts and activation
of BCL-2
2001

Copyright (c) 2001 INIST-CNRS. All rights reserved.

4/6/20 (Item 2 from file: 144)
13143960 PASCAL No.: 97-0403730
Establishment of a monkey kidney **epithelial** cell line with the
BARF1 open reading frame from Epstein-Barr virus
1997

Copyright (c) 1997 INIST-CNRS. All rights reserved. .

4/6/21 (Item 1 from file: 155)
13115134 21983099 PMID: 11986715
Malignant transformation of monkey kidney **epithelial** cell induced
by EBV **BARF1** gene and TPA]
Dec 2001

4/6/22 (Item 1 from file: 399)
DIALOG(R)File 399:(c) 2002 AMERICAN CHEMICAL SOCIETY. All rts. reserv.

Malignant transformation and immortalization assays in animal cells
transfected with the BARF1 gene
? t s4/7/8-11, 13, 14, 17
>>>Format 7 is not valid in file 143

4/7/8 (Item 8 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

11708757 BIOSIS NO.: 199800490488
Nucleic acid sequence-based amplification, a new method for analysis of
spliced and unspliced Epstein-Barr virus latent transcripts, and its
comparison with reverse transcriptase PCR.
AUTHOR: Brink Antoinette A T P; Vervoort Marcel B H J; Middeldorp Jaap M;
Meijer Chris J L M; Van Den Brule Adriaan J C(a)
AUTHOR ADDRESS: (a)Sect. Mol. Pathol., Dep. Univ. Hosp. Vrije Univ., P.O.
Box 7057, 1007 MB Amsterdam**Netherlands
JOURNAL: Journal of Clinical Microbiology 36 (11):p3164-3169 Nov., 1998
ISSN: 0095-1137
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Nucleic acid sequence-based amplification (NASBA) assays were
developed for direct detection of Epstein-Barr virus (EBV) transcripts
encoding EBV nuclear antigen 1 (EBNA1), latent membrane proteins (LMP) 1
and 2, and BamHIA rightward frame 1 (**BARF1**) and for the noncoding
EBV early RNA 1 (EBER1). The sensitivities of all NASBAs were at least

100 copies of specific in vitro-generated RNA. Furthermore, 1 EBV-positive JY cell in a background of 50,000 EBV-negative Ramos cells (the relative sensitivity) was detected by using the EBNA1, LMP1, and LMP2 NASBA assays. The relative sensitivity of the EBER1 NASBA was 100 EBV-positive cells, which was probably related to the loss of small RNA molecules during the isolation. The **BARF1** and LMP2 NASBAs were evaluated on clinical material. **BARF1** expression was found in 6 of 7 **nasopharyngeal** carcinomas (NPC) but in 0 of 22 Hodgkin's disease (HD) cases, whereas LMP2 expression was found in 7 of 7 NPCs and in 17 of 22 HD cases. For detection of EBNA1 transcripts in HLs (n = 12) and T- and B-cell non-Hodgkin's lymphomas (n = 3 and n = 2, respectively), NASBA was compared with reverse transcriptase (RT) PCR. Two samples were positive only with NASBA, and two other samples were positive only with RT-PCR; for all other samples, the RT-PCR and NASBA results were in agreement. We conclude that NASBA is suitable for sensitive and specific detection of the above-mentioned EBV transcripts, regardless of their splicing patterns and the presence of EBV DNA. The EBNA1, LMP2, and **BARF1** NASBAs developed in this study proved to be reliable assays for detection of the corresponding transcripts in EBV-positive clinical material.

4/7/9 (Item 9 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

11295044 BIOSIS NO.: 199800076376
Expression of the protein encoded by Epstein-Barr virus (EBV) **BARF1** open reading frame from a recombinant adenovirus system.
AUTHOR: de Turenne-Tesser M(a); Jolicoeur P; Ooka T
AUTHOR ADDRESS: (a)Virol. Mol., UMR 5537 CNRS, Fac. Med., RTH Laennec, rue G. Paradin, 69372 Lyon Cedex 08**France
JOURNAL: Virus Research 52 (1):p73-85 Nov., 1997
ISSN: 0168-1702
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Epstein-Barr virus (EBV) has been associated with human cancers of lymphocytic or **epithelial** origin, but viral functions implied in oncogenesis are not yet clear. We previously reported the oncogenic transformation of rodent fibroblast and human B lymphocyte cell lines by the **BARF1** coding sequence from EBV. We more recently observed immortalizing effects of this gene on monkey kidney primary **epithelial** cells. Here we describe an efficient recombinant adenovirus expression system which allowed us to characterize **BARF1** translation products, with the help of rabbit polyclonal antibodies raised to the entire protein. The present data demonstrate that **BARF1** encodes a 31-33 kDa hydrophobic protein, linked to cell membranes though also recovered in the cytosol, and recognized by human sera from patients with various EBV-related pathologies.

4/7/10 (Item 10 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

11025142 BIOSIS NO.: 199799646287
Establishment of a monkey kidney **epithelial** cell line with the **BARF1** open reading frame from Epstein-Barr virus.
AUTHOR: Wei Ming Xin; Turenne-Tessier Mireille De; Decaussin Gisele; Benet Gerard; Ooka Tadamasa(a)

AUTHOR ADDRESS: (a)Lab. Virol. Mol., IVMC, UMR 5537 CNRS, Fac. Med. R,
Laennec, Rue Guillaume Paradin, 69372 Lyon C**France
JOURNAL: Oncogene 14 (25):p3073-3081 1997
ISSN: 0950-9232
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: We previously reported that the **BARF1** (BamHI-A right frame 1) gene product from Epstein-Barr Virus (EBV) may have oncogenic properties since injection into new-born rats of transfected cell lines resulted in the development of **BARF1** expressing tumors, which were aggressive in the case of murine fibroblasts and transient in that of human B lymphocytes. As EBV has been associated with **nasopharyngeal carcinoma (NPC)** and evidence of **BARF1** transcription in this cancer was emerging from our biopsy analyses, we examined the effects of **BARF1** transfection into primate primary **epithelial** cells. The expression of the **BARF1** open reading frame in primary monkey kidney **epithelial** cells led us to the establishment of continuously dividing lines. The **BARF1** transfectants showed the major characteristics of immortalized cells: morphological change, short cell doubling time, ability to divide at low cell density and continuous growth over 50 passages. Injection of **BARF1** transfectants into nude mice did not induce any tumor. Established subclones were shown to be **epithelial** cells expressing known keratins as well as the **BARF1** coded mRNA and protein. This is the first report indicating that expression of the **BARF1** gene product in primary **epithelial** cells may contribute to the establishment of cell lines.

4/7/11 (Item 11 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

10745514 BIOSIS NO.: 199799366659
Antibody and antibody-dependent cellular cytotoxicity responses against the BamHI A rightward open-reading frame-1 protein of Epstein-Barr virus (EBV) in EBV-associated disorders.
AUTHOR: Tanner J E; Wei M X; Alfieri C; Ahmad A; Taylor P; Ooka T; Menezes J(a)
AUTHOR ADDRESS: (a)Lab. Immunoviol., Pediatric Res. Cent., Ste-Justine Hosp., 3175 Cote-Ste-Catherine, Montreal, Q**Canada
JOURNAL: Journal of Infectious Diseases 175 (1):p38-46 1997
ISSN: 0022-1899
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Antibody-dependent cellular cytotoxicity (ADCC) is an important antiviral effector mechanism. ADCC to the protein encoded by the Epstein-Barr virus (EBV) BamHI A rightward open-reading frame-1 (**BARF1**) was studied by transducing Raji-tk- cells with the **BARF1** gene using a retroviral expression vector. The transduced Raji cells expressed **BARF1** on the cell surface, as determined by flow cytometry. Sera from chronic and acute infectious mononucleosis and **nasopharyngeal carcinoma** patients were found to contain antibodies that react with the **BARF1** protein. When **BARF1**-expressing Raji cells were used as targets for ADCC, sera from several **nasopharyngeal carcinoma** patients demonstrated significant ADCC reactivity, whereas sera from healthy EBV-seronegative and -seropositive persons lacked such reactivity. **BARF1**-specific ADCC activity could be competitively inhibited with recombinant **BARF1** protein. The level of anti-**BARF1** antibody activity in sera of patients with EBV-associated diseases suggests that the **BARF1** protein may serve

as a target on EBV-infected cells for ADCC.

4/7/13 (Item 2 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

06651219 Genuine Article#: ZH268 Number of References: 60
Title: The Epstein-Barr virus **BARF1** gene encodes a novel, soluble
colony-stimulating factor-1 receptor
Author(s): Strockbine LD; Cohen JI; Farrah T; Lyman SD; Wagener F; DuBose
RF; Armitage RJ; Spriggs MK (REPRINT)
Corporate Source: IMMUNEX RES & DEV CORP, MOL BIOL, 51 UNIV
ST/SEATTLE//WA/98101 (REPRINT); IMMUNEX RES & DEV CORP, MOL
BIOL/SEATTLE//WA/98101; NIAID, CLIN INVEST LAB, NIH/BETHESDA//MD/20892
Journal: JOURNAL OF VIROLOGY, 1998, V72, N5 (MAY), P4015-4021
ISSN: 0022-538X Publication date: 19980500
Publisher: AMER SOC MICROBIOLOGY, 1325 MASSACHUSETTS AVENUE, NW,
WASHINGTON, DC 20005-4171
Language: English Document Type: ARTICLE
Abstract: Epstein-Barr virus (EBV) is a ubiquitous herpesvirus associated
with infectious mononucleosis and several tumors. The **BARF1** gene
is transcribed early after EBV infection from the BamHI A fragment of
the EBV genome. Evidence shown here indicates that the **BARF1**
protein is secreted into the medium of transfected cells and from
EBV-carrying B cells induced to allow lytic replication of the virus.
Expression cloning identified colony-stimulating factor-1 (CSF-1) as a
ligand for **BARF1**, Computer-assisted analyses indicated that
subtle amino acid sequence homology exists between **BARF1** and
c-fms, the cellular proto-oncogene that is the receptor for CSF-1.
Recombinant **BARF1** protein was found to be biologically active,
and it neutralized the proliferative effects of human CSF-1 in a
dose-dependent fashion when assayed in vitro. Since CSF-1 is a
pleiotropic cytokine best known for its differentiating effects on
macrophages, these data suggest that **BARF1** may function to
modulate the host immune response to EBV infection.

4/7/14 (Item 3 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

03058166 Genuine Article#: ND238 Number of References: 31
Title: EXPRESSION AND TUMORIGENICITY OF THE EPSTEIN-BARR-VIRUS **BARF1**
GENE IN HUMAN LOUCKES B-LYMPHOCYTE CELL-LINE
Author(s): WEI MX; MOULIN JC; DECAUSSIN G; BERGER F; OOKA T
Corporate Source: UNIV CLAUDE BERNARD, FAC MED ALEXIS CARREL, CNRS, IVMC, VIROL
MOLEC LAB, RUE G PARADIN/F-69372 LYON 08//FRANCE/; UNIV CLAUDE
BERNARD, FAC MED ALEXIS CARREL, CNRS, IVMC, VIROL MOLEC LAB, RUE G
PARADIN/F-69372 LYON 08//FRANCE/; PASTEUR MERIEUX/F-69280 MARCY
LETOILE//FRANCE/; HOP EDOUARD HERRIOT, SERV ANATOMOPATHOL/F-69374 LYON
08//FRANCE/
Journal: CANCER RESEARCH, 1994, V54, N7 (APR 1), P1843-1848
ISSN: 0008-5472
Language: ENGLISH Document Type: ARTICLE
Abstract: We previously showed that the Epstein-Barr virus, which encodes
the **BARF1** gene, could transform rodent fibroblasts. In this work,
the expression of the **BARF1** gene was studied in the human Louckes
B-lymphocyte cell line. Introduction of the **BARF1** open reading
frame under the control of the Mo-MuLV LTR promotor into nontumorigenic
Louckes lymphoid cells led to the activation of the c-myc protooncogene
and increased expression of the B-cell surface proteins, the

2/7/2 (Item 2 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

12522429 BIOSIS NO.: 200000275931

Unique transcription pattern of Epstein-Barr virus (EBV) in EBV-carrying gastric adenocarcinomas: Expression of the transforming **BARF1** gene.

AUTHOR: zur Hausen Axel; Brink Antoinette A T P; Craanen Mikael E;

Middeldorp Jaap M; Meijer Chris J L M; van den Brule Adriaan J C(a)

AUTHOR ADDRESS: (a)Department of Pathology, Section Molecular Pathology, University Hospital Vrije Universiteit, 1007 MB, Amsterdam**Netherlands

JOURNAL: Cancer Research 60 (10):p2745-2748 May 15, 2000

MEDIUM: print.

ISSN: 0008-5472

DOCUMENT TYPE: Article

RECORD TYPE: Abstract

LANGUAGE: English

SUMMARY LANGUAGE: English

ABSTRACT: Approximately 10% of gastric adenocarcinomas worldwide are associated with human EBV. These carcinomas generally do not express the latent membrane protein 1 (LMP1), the major known EBV oncogene. Recently, another EBV gene (i.e., **BARF1** (BamHI A rightward open reading frame)) was shown to have transforming and immortalizing capacities. Therefore, in this study, we investigated the expression of **BARF1** in EBV-carrying gastric adenocarcinomas in relation to the expression of other latent EBV transcripts. In the present study, 10 of 132 gastric adenocarcinomas tested positive for EBV using EBER1/2-RNA in situ hybridization. We demonstrate **BARF1** gene transcription in nine EBV-carrying gastric adenocarcinomas (with sufficient RNA quality) using the **BARF1**-specific nucleic acid sequence-based amplification assay. In addition, we also detected other latent EBV transcripts (i.e., **BARF0**-, **LMP2A**-, and **Q/K-driven EBNA1** transcripts in these carcinomas using reverse transcription-PCR analysis. No expression of **LMP1**, **EBNA2**, and **ZEBRA** (either at transcription or protein level) was found. In addition, two cases were positive for **BHRF1** transcripts, the viral bcl-2 homologue. Thus, together with **BARF1** transcription, a unique and distinct EBV latency type has been found in EBV-associated gastric adenocarcinomas. Because **BARF1** exerts immortalizing effects on human epithelial cells in vitro and EBV-carrying gastric adenocarcinomas lack the expression of **LMP1**, the **BARF1** gene might act as the viral oncogene in EBV-carrying gastric carcinomas. The **BARF1** gene offers an alternative way for EBV-mediated oncogenesis other than **LMP1**.

2/7/3 (Item 3 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

12511243 BIOSIS NO.: 200000264745

Expression of the transforming Epstein-Barr virus **BARF1** gene in EBV-carrying gastric adenocarcinomas.

AUTHOR: van den Brule Adriaan J(a); zur Hausen Axel; Brink Antoinette A;

Craanen Mikael E; Middeldorp Jaap M; Meijer Chris J

AUTHOR ADDRESS: (a)Dept of Pathology, Free Univ Amsterdam, Amsterdam** Netherlands

JOURNAL: Gastroenterology 118 (4 Suppl. 2 Part 1):pA60 April, 2000

MEDIUM: print.

CONFERENCE/MEETING: 101st Annual Meeting of the American Gastroenterological Association and the Digestive Disease Week. San Diego, California, USA May 21-24, 2000

SPONSOR: American Gastroenterological Association

ISSN: 0016-5085
RECORD TYPE: Citation
LANGUAGE: English
SUMMARY LANGUAGE: English

2/7/4 (Item 4 from file: 5)
DIALOG(R)File 5:Biosis Previews(R)
(c) 2002 BIOSIS. All rts. reserv.

11708757 BIOSIS NO.: 199800490488
Nucleic acid sequence-based amplification, a new method for analysis of spliced and unspliced Epstein-Barr virus latent transcripts, and its comparison with reverse transcriptase PCR.
AUTHOR: Brink Antoinette A T P; Vervoort Marcel B H J; Middeldorp Jaap M; Meijer Chris J L M; Van Den Brule Adriaan J C(a)
AUTHOR ADDRESS: (a)Sect. Mol. Pathol., Dep. Univ. Hosp. Vrije Univ., P.O. Box 7057, 1007 MB Amsterdam**Netherlands
JOURNAL: Journal of Clinical Microbiology 36 (11):p3164-3169 Nov., 1998
ISSN: 0095-1137
DOCUMENT TYPE: Article
RECORD TYPE: Abstract
LANGUAGE: English

ABSTRACT: Nucleic acid sequence-based amplification (NASBA) assays were developed for direct detection of Epstein-Barr virus (EBV) transcripts encoding EBV nuclear antigen 1 (**EBNA1**), latent membrane proteins (LMP) 1 and 2, and BamH1A rightward frame 1 (**BARF1**) and for the noncoding EBV early RNA 1 (EBER1). The sensitivities of all NASBAs were at least 100 copies of specific in vitro-generated RNA. Furthermore, 1 EBV-positive JY cell in a background of 50,000 EBV-negative Ramos cells (the relative sensitivity) was detected by using the **EBNA1**, LMP1, and LMP2 NASBA assays. The relative sensitivity of the EBER1 NASBA was 100 EBV-positive cells, which was probably related to the loss of small RNA molecules during the isolation. The **BARF1** and LMP2 NASBAs were evaluated on clinical material. **BARF1** expression was found in 6 of 7 nasopharyngeal carcinomas (NPC) but in 0 of 22 Hodgkin's disease (HD) cases, whereas LMP2 expression was found in 7 of 7 NPCs and in 17 of 22 HD cases. For detection of **EBNA1** transcripts in HLs (n = 12) and T- and B-cell non-Hodgkin's lymphomas (n = 3 and n = 2, respectively), NASBA was compared with reverse transcriptase (RT) PCR. Two samples were positive only with NASBA, and two other samples were positive only with RT-PCR; for all other samples, the RT-PCR and NASBA results were in agreement. We conclude that NASBA is suitable for sensitive and specific detection of the above-mentioned EBV transcripts, regardless of their splicing patterns and the presence of EBV DNA. The **EBNA1**, LMP2, and **BARF1** NASBAs developed in this study proved to be reliable assays for detection of the corresponding transcripts in EBV-positive clinical material.

2/7/5 (Item 1 from file: 34)
DIALOG(R)File 34:SciSearch(R) Cited Ref Sci
(c) 2002 Inst for Sci Info. All rts. reserv.

08668991 Genuine Article#: 314MJ Number of References: 34
Title: Unique transcription pattern of Epstein-Barr virus (EBV) in EBV-carrying gastric adenocarcinomas: Expression of the transforming **BARF1** gene
Author(s): zurHausen A; Brink AATP; Craanen ME; Middeldorp JM; Meijer CJLM; vandenBrule AJC (REPRINT)
Corporate Source: FREE UNIV AMSTERDAM HOSP, DEPT PATHOL, SECT MOL PATHOL,

POB 7057/NL-1007 MB AMSTERDAM//NETHERLANDS/ (REPRINT); FREE UNIV
AMSTERDAM HOSP, DEPT PATHOL, SECT MOL PATHOL/NL-1007 MB
AMSTERDAM//NETHERLANDS//; FREE UNIV AMSTERDAM HOSP, DEPT
GASTROENTEROL/NL-1007 MB AMSTERDAM//NETHERLANDS/
Journal: CANCER RESEARCH, 2000, V60, N10 (MAY 15), P2745-2748
ISSN: 0008-5472 Publication date: 20000515
Publisher: AMER ASSOC CANCER RESEARCH, PO BOX 11806, BIRMINGHAM, AL 35202
Language: English Document Type: ARTICLE
Abstract: Approximately 10% of gastric adenocarcinomas worldwide are
associated with human EBV, These carcinomas generally do not express
the latent membrane protein 1 (LMP1), the major known EBV oncogene,

Recently, another EBV gene [i.e., **BARF1** (BamHI A rightward
open reading frame)] was shown to have transforming and immortalizing
capacities, Therefore, in this study, we investigated the expression of
BARF1 in EBV-carrying gastric adenocarcinomas in relation to the
expression of other latent EBV transcripts.

In the present study, 10 of 132 gastric adenocarcinomas tested
positive for EBV using EBER1/2-RNA in situ hybridization, We
demonstrate **BARF1** gene transcription in nine EBV-carrying gastric
adenocarcinomas (with sufficient RNA quality) using the **BARF1**
-specific nucleic acid sequence-based amplification assay. In addition,
we also detected other latent EBV transcripts (i.e., **BARF0**-, **LMP2A**-,
and **Q/K**-driven **EBNA1** transcripts in these carcinomas using
reverse transcription-PCR analysis. No expression of **LMP1**, **EBNA2**, and
ZEBRA (either at transcription or protein level) was found. In
addition, two cases were positive for **BHRF1** transcripts, the viral
bcl-2 homologue, Thus, together with **BARF1** transcription, a
unique and distinct EBV latency type has been found in EBV-associated
gastric adenocarcinomas.

Because **BARF1** exerts immortalizing effects on human
epithelial cells in vitro and EBV-carrying gastric adenocarcinomas lack
the expression of **LMP1** the **BARF1** gene might act as the viral
oncogene in EBV-carrying gastric carcinomas. The **BARF1** gene
offers an alternative way for EBV-mediated oncogenesis other than **LMP1**.

2/7/6 (Item 1 from file: 76)
DIALOG(R) File 76:Life Sciences Collection
(c) 2002 Cambridge Sci Abs. All rts. reserv.

02492243 4737903

Unique Transcription Pattern of Epstein-Barr Virus (EBV) in EBV-carrying
Gastric Adenocarcinomas: Expression of the Transforming **BARF1** Gene
Hausen, A.Z.; Brink, A.A.T.P.; Craanen, M.E.; Middeldorp, J.M.; Meijer,
C.J.L.M.; van den Brule, A.J.C.

Department of Pathology, Section Molecular Pathology, University Hospital
Vrije Universiteit, P.O. Box 7057, 1007 MB Amsterdam, the Netherlands
Cancer Research vol. 60, no. 10, pp. 2745-2748 (2000)
ISSN: 0008-5472

DOCUMENT TYPE: Journal article LANGUAGE: ENGLISH
SUBFILE: Virology & AIDS Abstracts

Approximately 10% of gastric adenocarcinomas worldwide are associated
with human EBV. These carcinomas generally do not express the latent
membrane protein 1 (**LMP1**), the major known EBV oncogene. Recently, another
EBV gene [i.e., **BARF1** (BamHI A rightward open reading frame)] was
shown to have transforming and immortalizing capacities. Therefore, in this
study, we investigated the expression of **BARF1** in EBV-carrying
gastric adenocarcinomas in relation to the expression of other latent EBV

transcripts. In the present study, 10 of 132 gastric adenocarcinomas tested positive for EBV using EBER1/2-RNA in situ hybridization. We demonstrate **BARF1** gene transcription in nine EBV-carrying gastric adenocarcinomas (with sufficient RNA quality) using the **BARF1**-specific nucleic acid sequence-based amplification assay. In addition, we also detected other latent EBV transcripts (i.e., **BARF0**-, **LMP2A**-, and **Q/K**-driven **EBNA1** transcripts in these carcinomas using reverse transcription-PCR analysis. No expression of **LMP1**, **EBNA2**, and **ZEBRA** (either at transcription or protein level) was found. In addition, two cases were positive for **BHRF1** transcripts, the viral **bcl-2** homologue. Thus, together with **BARF1** transcription, a unique and distinct EBV latency type has been found in EBV-associated gastric adenocarcinomas. Because **BARF1** exerts immortalizing effects on human epithelial cells in vitro and EBV-carrying gastric adenocarcinomas lack the expression of **LMP1**, the **BARF1** gene might act as the viral oncogene in EBV-carrying gastric carcinomas. The **BARF1** gene offers an alternative way for EBV-mediated oncogenesis other than **LMP1**.

2/7/7 (Item 1 from file: 98)
DIALOG(R)File 98:General Sci Abs/Full-Text
(c) 2002 The HW Wilson Co. All rts. reserv.

04273985 H.W. WILSON RECORD NUMBER: BGSA00023985 (THIS IS THE FULLTEXT)
Role of cytotoxic T lymphocytes in Epstein-Barr virus-associated diseases.
Khanna, Rajiv
Burrows, Scott R
Annual Review of Microbiology v. 54 (2000), p. 19-48
LANGUAGE: English
COUNTRY OF PUBLICATION: United States
WORD COUNT: 11989

ABSTRACT: Adaptation of persistent infection within the cells of the immune system is a unique characteristic of gamma herpes viruses. A classic example of this is Epstein-Barr virus (EBV), which may have co-evolved with *Homo sapiens* over millions of years, thus achieving a balance between viral persistence and immune control. In this review, we present an overview of virus and the host immune system interactions that regulate the life-long host-virus relationship in healthy virus carriers and EBV-associated diseases. Extensive analysis of cytotoxic T lymphocyte-mediated immune responses in healthy virus carriers has revealed unique mechanisms used by EBV to maintain a benign persistent state in vivo. On the other hand, this relationship in EBV-associated diseases favors the escape of the virus from the hostile effects of the immune response. This escape is achieved by either down-regulating the expression of highly immunogenic antigens of the virus or by direct modulation of the host cytotoxic T lymphocyte response by virus-encoded proteins. Reprinted by permission of the publisher.

TEXT:

Key Words latency, lymphoma, immunomodulation, epitope, immune escape, immunotherapy

INTRODUCTION

Epstein-Barr virus (EBV) is the best known and most widely studied herpes virus, owing to its clinical and oncogenic importance. Although EBV was the first virus implicated in a human cancer, its ubiquitous prevalence in diverse human populations made it difficult to firmly establish the link between this virus and lymphomas. In early 1964, Epstein and colleagues reported that they could continuously culture cells from a Burkitt's lymphoma (BL) patient, and, using electron microscopy, they identified a herpes viral particle (19). This finding was the culmination of years of epidemiological, cell biology, and electron microscopic studies. In fact,

transferrin receptor, CD21, and CD23. **BARF1**-expressing cells induced a diffuse lymphoma-like tumor in newborn rats treated with anti-thymocyte serum that was, however, transient and regressed after 3-4 weeks as the immune system recovered. The tumor induction was similar to that observed with lymphoid cell lines in vitro generated by infection with the B95-8 virus strain, in which lytic antigens are expressed at low levels. After long-term culture, Louckes cell clones lost expression of the **BARF1** gene and were unable to induce tumors.

4/7/17 (Item 2 from file: 76)
DIALOG(R)File 76:Life Sciences Collection
(c) 2002 Cambridge Sci Abs. All rts. reserv.

02217680 4231116

Expression of the protein encoded by Epstein-Barr virus (EBV) **BARF1**
open reading frame from a recombinant adenovirus system

de Turenne Tessier, M.; Jolicoeur, P.; Ooka, T.

Virologie Molculaire, UMR 5537 CNRS, Facultede Medecine RTH Laennec, rue
G. Paradin 69372, Lyon Cedex 08, France

VIRUS RES. vol. 52, no. 1, pp. 73-85 (1997)

ISSN: 0168-1702

DOCUMENT TYPE: Journal article LANGUAGE: ENGLISH

SUBFILE: Virology & AIDS Abstracts

Epstein-Barr virus (EBV) has been associated with human cancers of lymphocytic or **epithelial** origin, but viral functions implied in oncogenesis are not yet clear. We previously reported the oncogenic transformation of rodent fibroblast and human B lymphocyte cell lines by the **BARF1** coding sequence from EBV. We more recently observed immortalizing effects of this gene on monkey kidney primary **epithelial** cells. Here we describe an efficient recombinant adenovirus expression system which allowed us to characterize **BARF1** translation products, with the help of rabbit polyclonal antibodies raised to the entire protein. The present data demonstrate that **BARF1** encodes a 31-33 kDa hydrophobic protein, linked to cell membranes though also recovered in the cytosol, and recognized by human sera from patients with various EBV-related pathologies.

?